FIG. 1

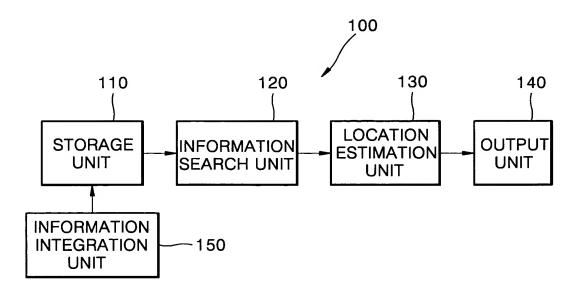


FIG. 3

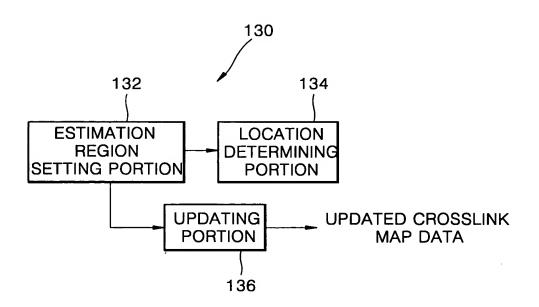


FIG. 2

. Orient	+	+	+	+	+	+	+	+
Annot . End	84190	82311	188	2	84189	82310	188	2
Annot . Start	-	1	1	-	-	1	1	1
Annot . ID	BRCA2.TRANS	BRCA2. CDS	BRCA2.EXON1	dbSNP :206118	BRCA2.TRANS	BRCA2. CDS	BRCA2.EXON1	dbSNP :206118
Annot . Type	TRANS	SOO	EXON	SNP	TRANS	SOO	NOXE	dNS
Chr	31045834	31044936	30961832	30961821	30960165	30959267	30876164	30876152
Chr Start	30961644	30962625	30961644	30961820	30875976	30876957	30875976	30876151
Chr	Chr13	Chr13	Chr13	Chr13	Chr13	Chr13	Chr13	Chr13
Genome Version	200104	200104	200104	200104	200206	200206	200206	200206
Genome	ncsc	OCSC	oson	OCSC	ncsc	ncsc	OCSC	OSON

FIG. 4

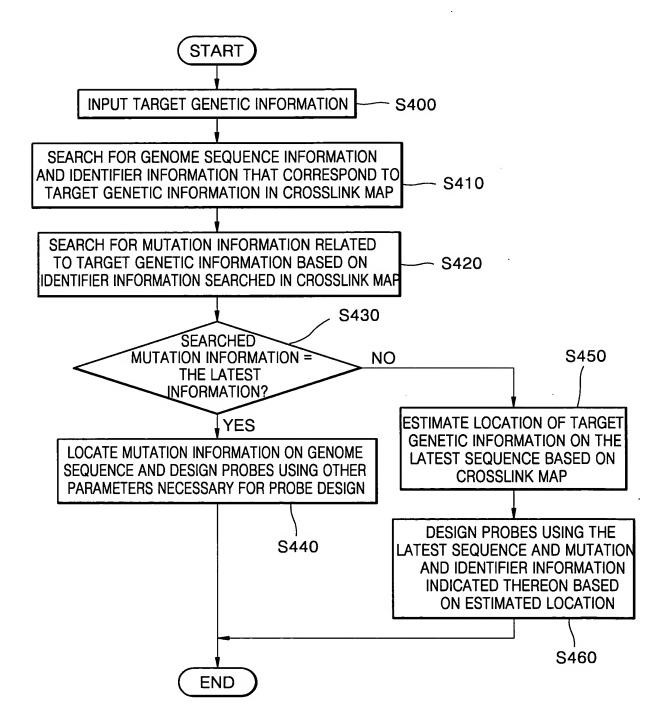


FIG. 5

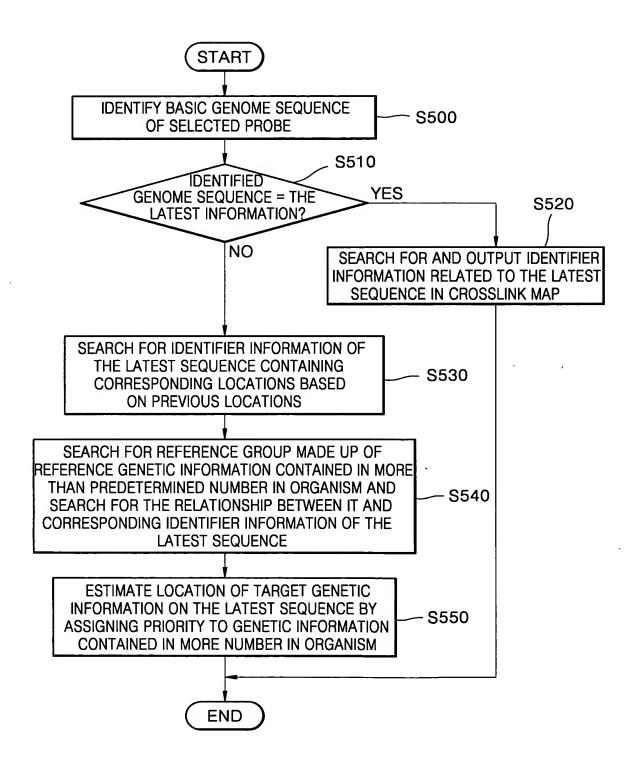


FIG. 6

TRANS: 30875976 - 30960165 (84189) CDS : 30876957 - 30959267 (82310) EXON : 30958658 - 30959707 (1049) EXON: 30876918 - 30877024 (106) EXON: 30955185 - 30955430 (245) EXON: 30875976 - 30876164 (188) EXON: 30957394 - 30957541 (147) SNPN: 30876151 - 30876152 UCSC.200206 SNPN: 31041006 - 31041007 [1799968] SNPN: 30961820 - 30961821 [206118] : 30962625 - 31044936 (82311) TRANS: 30961644 - 31045834 (84190) EXON: 30961644 - 30961832 (188) EXON: 31043063 - 31043210 (147) EXON: 31025908 - 31026072 (164) EXON: 31040851 - 31041096 (245) UCSC.200104

206118

GCCTCATATGTTAATTGCTGCAAGCAACCTCCAGTGGCGAC **TAATTGCTGCAAGCAACCTC** GCCTCATATGTTGCTGCAAGCAACCTCCAGTGGCGAC TAATTACTGCAAGCAACCTC

31041020

31041000

30955345 30955325